



## SEQUENCE LISTING

Flint, Andrew J.  
Cool, Deborah E.

<1> IMPROVED ASSAY FOR PROTEIN TYROSINE  
KINASES

<110> 200125.401

<111> US/09/788,626

<112> 2001-02-13

<160> 40

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<221> VARIANT

<222> (1)...(1)

<223> Xaa = Ile or Val

<221> VARIANT

<222> (4)...(4)

<223> Xaa = any amino acid

<221> VARIANT

<222> (7)...(7)

<223> Xaa = any amino acid

<221> VARIANT

<222> (8)...(8)

<223> Xaa = any amino acid

<221> VARIANT

<222> (10)...(10)

<223> Xaa = Ser or Thr

<224> Unique signature sequence motif which is invariant  
among all PTPs.

<210> 1

Gla His Cys Xaa Ala Gly Xaa Xaa Arg Xaa Gly

1

5

10

<210> 2

<211> 254

<212> PRT

<213> Homo sapiens

<400> 2

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Asp Phe Pro Cys Arg Val Ala Lys Leu Pro Lys Asn Lys Asn Arg Asn
 1           5           10           15
Arg Tyr Arg Asp Val Ser Pro Phe Asp His Ser Arg Ile Lys Leu His
 20           25           30
Gln Glu Asp Asn Asp Tyr Ile Asn Ala Ser Leu Ile Lys Met Glu Glu
 35           40           45
Ala Gln Arg Ser Tyr Ile Leu Thr Gln Gly Pro Leu Pro Asn Thr Cys
 50           55           60
Gly His Phe Trp Glu Met Val Trp Glu Gln Lys Ser Arg Gly Val Val
 65           70           75           80
Met Leu Asn Arg Val Met Glu Lys Gly Ser Leu Lys Cys Ala Gln Tyr
 85           90           95
Trp Pro Gln Lys Glu Glu Lys Glu Met Ile Phe Glu Asp Thr Asn Leu
100          105          110
Lys Leu Thr Leu Ile Ser Glu Asp Ile Lys Ser Tyr Tyr Thr Val Leu
115          120          125
Glu Leu Glu Asn Leu Thr Thr Gln Glu Thr Arg Glu Ile Leu His Phe
130          135          140
His Tyr Thr Thr Trp Pro Asp Phe Gly Val Pro Glu Ser Pro Ala Ser
145          150          155          160
Phe Leu Asn Phe Leu Phe Lys Val Arg Glu Ser Gly Ser Leu Ser Pro
165          170          175
Glu His Gly Pro Val Val Val His Cys Ser Ala Gly Ile Gly Arg Ser
180          185          190
Gly Thr Phe Cys Leu Ala Asp Thr Cys Leu Leu Leu Met Asp Lys Arg
195          200          205
Lys Asp Pro Ser Ser Val Asp Ile Lys Lys Val Leu Leu Glu Met Arg
210          215          220
Lys Phe Arg Met Gly Leu Ile Gln Thr Ala Asp Gln Leu Arg Phe Ser
225          230          235          240
Tyr Leu Ala Val Ile Glu Gly Ala Lys Phe Ile Met Gly Asp
245          250

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<210> 3

<211> 251

<212> PRT

<213> Homo sapiens

<400> 3

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Asp Tyr Pro His Arg Val Ala Lys Phe Pro Glu Asn Arg Asn Arg Asn
 1           5           10           15
Arg Tyr Arg Asp Val Ser Pro Tyr Asp His Ser Arg Val Leu Gln Asn
 20           25           30
Ala Glu Asn Asp Tyr Ile Asn Ala Ser Leu Val Asp Ile Glu Glu Ala
 35           40           45
Gln Arg Ser Tyr Ile Leu Thr Gln Gly Pro Leu Pro Asn Thr Cys Cys
 50           55           60
His Phe Trp Leu Met Val Trp Gln Gln Lys Thr Lys Ala Val Val Met
 65           70           75           80
Leu Asn Arg Ile Val Glu Lys Glu Ser Val Lys Cys Ala Gln Tyr Trp
 85           90           95

```

Pro Thr Asp Asp Gln Glu Met Leu Phe Lys Glu Thr Gly Phe Ser Val  
 100 105 110  
 Lys Leu Leu Ser Glu Asp Val Lys Ser Tyr Tyr Thr Val Leu Gln Leu  
 115 120 125  
 Glu Asn Ile Asn Ser Gly Glu Thr Arg Thr Ile Ser His Phe His Tyr  
 130 135 140  
 Thr Thr Trp Pro Asp Phe Gly Val Pro Glu Ser Pro Ala Ser Phe Leu  
 145 150 155 160  
 Asn Phe Leu Phe Lys Val Arg Glu Ser Gly Ser Leu Asn Pro Asp His  
 165 170 175  
 Gly Pro Ala Val Ile His Cys Ser Ala Gly Ile Gly Arg Ser Gly Thr  
 180 185 190  
 Phe Ser Leu Val Asp Thr Cys Leu Val Leu Met Glu Lys Gly Asp Asp  
 195 200 205  
 Ile Asn Ile Lys Gln Val Leu Leu Asn Met Arg Lys Tyr Arg Met Gly  
 210 215 220  
 Leu Ile Gln Thr Pro Asp Gln Leu Arg Phe Ser Tyr Met Ala Ile Ile  
 225 230 235 240  
 Glu Gly Ala Lys Cys Ile Lys Gly Asp Ser Ser  
 245 250

<210> 4  
 <211> 317  
 <212> PRT  
 <213> Homo sapiens

<400> 4  
 Gly Ile Thr Ala Asp Ser Ser Asn His Pro Asp Asn Lys His Lys Asn  
 1 5 10 15  
 Arg Tyr Ile Asn Ile Val Ala Tyr Asp His Ser Arg Val Lys Leu Ala  
 20 25 30  
 Gln Leu Ala Glu Lys Asp Gly Lys Leu Thr Asp Tyr Ile Asn Ala Asn  
 35 40 45  
 Tyr Val Asp Gly Tyr Asn Arg Pro Lys Ala Tyr Ile Ala Ala Gln Gly  
 50 55 60  
 Pro Leu Lys Ser Thr Ala Glu Asp Phe Trp Arg Met Ile Trp Glu His  
 65 70 75 80  
 Asn Val Glu Val Ile Val Met Ile Thr Asn Leu Val Glu Lys Gly Arg  
 85 90 95  
 Arg Lys Cys Asp Gln Tyr Trp Pro Pro Ala Asp Gly Ser Glu Glu Tyr  
 100 105 110  
 Gly Asn Phe Leu Val Thr Gln Lys Ser Val Gln Val Leu Ala Tyr Tyr  
 115 120 125  
 Thr Val Phe Thr Leu Arg Asn Thr Lys Ile Lys Lys Gly Ser Gln Lys  
 130 135 140  
 Gly Arg Pro Ser Gly Arg Val Val Thr Gln Tyr His Tyr Thr Gln Trp  
 145 150 155 160  
 Pro Asp Met Gly Val Pro Glu Tyr Ser Leu Pro Val Leu Thr Phe Val  
 165 170 175  
 Arg Lys Ala Ala Tyr Ala Lys Arg His Ala Val Gly Pro Val Val Val  
 180 185 190  
 His Cys Ser Ala Gly Val Gly Arg Thr Gly Thr Tyr Ile Val Leu Asp  
 195 200 205  
 Ser Met Leu Gln Gln Ile Gln His Glu Gly Thr Val Asn Ile Phe Gly

210 215 220  
 Phe Leu Lys His Ile Arg Ser Gln Arg Asn Tyr Leu Val Gln Thr Glu  
 225 230 235 240  
 Glu Gln Tyr Val Phe Ile His Asp Thr Leu Val Glu Ala Ile Leu Ser  
 245 250 255  
 Lys Glu Thr Glu Val Val Leu Asp Ser Met Leu Gln Gln Ile Gln His  
 260 265 270  
 Glu Gly Thr Val Asn Ile Phe Gly Phe Leu Lys His Ile Arg Ser Gln  
 275 280 285  
 Arg Asn Tyr Leu Val Gln Thr Glu Glu Gln Tyr Val Phe Ile His Asp  
 290 295 300  
 Thr Leu Val Glu Ala Ile Leu Ser Lys Glu Thr Glu Val  
 305 310 315

<210> 5

<211> 316

<212> PPT

<213> Homo sapiens

<400> 5

Gly Ile Thr Ala Asp Ser Ser Asn His Pro Asp Asn Lys His Lys Asn  
 1 5 10 15  
 Arg Tyr Ile Asn Ile Val Ala Tyr Asp His Ser Arg Val Lys Leu Ala  
 20 25 30  
 Gln Leu Ala Glu Lys Asp Gly Lys Leu Thr Asp Tyr Ile Asn Ala Asn  
 35 40 45  
 Tyr Val Asp Gly Tyr Asn Arg Pro Lys Ala Tyr Ile Ala Ala Gln Gly  
 50 55 60  
 Pro Leu Lys Ser Thr Ala Glu Asp Phe Trp Arg Met Ile Trp Glu His  
 65 70 75 80  
 Asn Val Glu Val Ile Val Met Ile Thr Asn Leu Val Glu Lys Gly Arg  
 85 90 95  
 Arg Lys Cys Asp Gln Tyr Trp Pro Ala Asp Gly Ser Glu Glu Tyr Gly  
 100 105 110  
 Asn Phe Leu Val Thr Gln Lys Ser Val Gln Val Leu Ala Tyr Tyr Thr  
 115 120 125  
 Val Phe Thr Leu Arg Asn Thr Lys Ile Lys Lys Gly Ser Gln Lys Gly  
 130 135 140  
 Arg Pro Ser Gly Arg Val Val Thr Gln Tyr His Tyr Thr Gln Trp Pro  
 145 150 155 160  
 Asp Met Gly Val Pro Glu Tyr Ser Leu Pro Val Leu Thr Phe Val Arg  
 165 170 175  
 Lys Ala Ala Tyr Ala Lys Arg His Ala Val Gly Pro Val Val Val His  
 180 185 190  
 Cys Ser Ala Gly Val Gly Arg Thr Gly Thr Tyr Ile Val Leu Asp Ser  
 195 200 205  
 Met Leu Gln Gln Ile Gln His Glu Gly Thr Val Asn Ile Phe Gly Phe  
 210 215 220  
 Leu Lys His Ile Arg Ser Gln Arg Asn Tyr Leu Val Gln Thr Glu Glu  
 225 230 235 240  
 Gln Tyr Val Phe Ile His Asp Thr Leu Val Glu Ala Ile Leu Ser Lys  
 245 250 255  
 Glu Thr Glu Val Val Leu Asp Ser Met Leu Gln Gln Ile Gln His Glu  
 260 265 270

Gly Thr Val Asn Ile Phe Gly Ile Leu Lys His Ile Arg Ser Gln Arg  
 275 280 285  
 Asn Tyr Leu Val Gln Thr Glu Gln Gln Tyr Val Phe Ile His Asp Thr  
 290 295 300  
 Leu Val Glu Ala Ile Leu Ser Lys Glu Thr Glu Val  
 305 310 315

<210> 6  
 <211> 319  
 <212> PRT  
 <213> Homo sapiens

<400> 6  
 Asn Ile Thr Ala Glu His Ser Asn His Pro Glu Asn Lys His Lys Asn  
 1 5 10 15  
 Arg Tyr Ile Asn Ile Leu Ala Tyr Asp His Ser Arg Val Lys Leu Arg  
 20 25 30  
 Pro Leu Pro Gly Lys Asp Ser Lys His Ser Asp Tyr Ile Asn Ala Asn  
 35 40 45  
 Tyr Val Asp Gly Tyr Asn Lys Ala Lys Ala Tyr Ile Ala Thr Gln Gly  
 50 55 60  
 Pro Leu Lys Ser Thr Phe Glu Asp Phe Trp Arg Met Ile Trp Glu Gln  
 65 70 75 80  
 Asn Thr Gly Ile Ile Val Met Ile Thr Asn Leu Val Glu Lys Gly Arg  
 85 90 95  
 Arg Lys Cys Asp Gln Tyr Trp Pro Thr Glu Asn Ser Glu Glu Tyr Gly  
 100 105 110  
 Asn Ile Ile Val Thr Leu Lys Ser Thr Lys Ile His Ala Cys Tyr Thr  
 115 120 125  
 Val Phe Ser Ile Arg Asn Thr Lys Val Lys Lys Gly Gln Lys Gly Asn  
 130 135 140  
 Pro Lys Gly Arg Gln Asn Glu Arg Val Val Ile Gln Tyr His Tyr Thr  
 145 150 155 160  
 Gln Trp Pro Asp Met Gly Val Pro Glu Tyr Ala Leu Pro Val Leu Thr  
 165 170 175  
 Phe Val Arg Arg Ser Ser Ala Ala Arg Met Pro Glu Thr Gly Pro Val  
 180 185 190  
 Leu Val His Cys Ser Ala Gly Val Gly Arg Thr Gly Thr Tyr Ile Val  
 195 200 205  
 Ile Asp Ser Met Leu Gln Gln Ile Lys Asp Lys Ser Thr Val Asn Val  
 210 215 220  
 Leu Gly Phe Leu Lys His Ile Arg Thr Gln Arg Asn Tyr Leu Val Gln  
 225 230 235 240  
 Thr Glu Glu Gln Tyr Ile Ile Ile His Asp Ala Leu Leu Glu Ala Ile  
 245 250 255  
 Leu Gly Lys Glu Thr Glu Val Val Ile Asp Ser Met Leu Gln Gln Ile  
 260 265 270  
 Lys Asp Lys Ser Thr Val Asn Val Leu Gly Phe Leu Lys His Ile Arg  
 275 280 285  
 Thr Gln Arg Asn Tyr Leu Val Glu Thr Glu Glu Glu Tyr Ile Phe Ile  
 290 295 300  
 His Asp Ala Leu Leu Glu Ala Ile Leu Gly Lys Glu Thr Glu Val  
 305 310 315

<210> 7  
 <211> 313  
 <212> PRT  
 <213> *Drosophila melanogaster*

<400> 7  
 Asp Leu Pro Cys Glu His Ser Gln His Pro Glu Asn Lys Arg Lys Asn  
 1 5 10 15  
 Arg Tyr Leu Asn Ile Thr Ala Tyr Asp His Ser Arg Val His Leu His  
 20 25 30  
 Pro Thr Pro Gly Gln Lys Lys Asn Leu Asp Tyr Ile Asn Ala Asn Phe  
 35 40 45  
 Ile Asp Gly Tyr Gln Lys Gly His Ala Phe Ile Gly Thr Gln Gly Pro  
 50 55 60  
 Leu Pro Asp Thr Phe Asp Cys Phe Trp Arg Met Ile Trp Glu Gln Arg  
 65 70 75 80  
 Val Ala Ile Ile Val Met Ile Thr Asn Leu Val Glu Arg Gly Arg Arg  
 85 90 95  
 Lys Cys Asp Met Tyr Trp Phe Lys Asp Gly Val His Thr Tyr Gly Val  
 100 105 110  
 Ile Gln Val Lys Leu Ile Glu Glu Glu Val Met Ser Thr Tyr Thr Val  
 115 120 125  
 Leu Gln Ile Lys His Leu Lys Leu Lys Lys Lys Lys Gln Cys Asn Thr  
 130 135 140  
 Glu Lys Leu Val Tyr Gln Tyr His Tyr Thr Asn Trp Phe Asp His Gly  
 145 150 155 160  
 Thr Pro Asp His Pro Leu Pro Val Leu Asn Phe Val Lys Lys Ser Ser  
 165 170 175  
 Ala Ala Asn Pro Ala Glu Ala Gly Phe Ile Val Val His Tyr Ser Ala  
 180 185 190  
 Gly Val Gly Arg Thr Gly Thr Tyr Ile Val Leu Asp Ala Met Leu Lys  
 195 200 205  
 Gln Ile Gln Gln Lys Asn Ile Val Asn Val Phe Gly Phe Leu Arg His  
 210 215 220  
 Ile Arg Ala Gln Arg Asn Phe Leu Val Gln Thr Glu Glu Gln Tyr Ile  
 225 230 235 240  
 Phe Leu His Asp Ala Leu Val Glu Ala Ile Ala Ser Gly Glu Thr Asn  
 245 250 255  
 Leu Val Leu Asp Ala Met Leu Lys Gln Ile Gln Gln Lys Asn Ile Val  
 260 265 270  
 Asn Val Phe Gly Phe Leu Arg His Ile Arg Ala Gln Arg Asn Phe Leu  
 275 280 285  
 Val Gln Thr Glu Glu Gln Tyr Ile Phe Leu His Asp Ala Leu Val Glu  
 290 295 300  
 Ala Ile Ala Ser Gly Glu Thr Asn Leu  
 305 310

<210> 8  
 <211> 306  
 <212> PRT  
 <213> *Homo sapiens*

<400> 8  
 Gln Phe Thr Trp Gln Asn Ser Asn Leu Gln Val Asn Lys Trp Lys Asn

1 5 10 15  
 Arg Tyr Ala Asn Val Ile Ala Tyr Asp His Ser Arg Val Ile Leu Thr  
 20 25 30  
 Ser Ile Asp Gly Val Pro Gly Ser Asp Tyr Ile Asn Ala Asn Tyr Ile  
 35 40 45  
 Asp Gly Tyr Arg Lys Gln Asn Ala Tyr Ile Ala Thr Gln Gly Pro Leu  
 50 55 60  
 Pro Glu Thr Met Gly Asp Phe Trp Arg Met Val Trp Glu Gln Arg Thr  
 65 70 75 80  
 Ala Thr Val Val Met Met Thr Arg Leu Glu Glu Lys Ser Arg Val Lys  
 85 90 95  
 Cys Asp Gln Tyr Trp Pro Ala Arg Gly Thr Glu Thr Cys Gly Leu Ile  
 100 105 110  
 Gln Val Thr Leu Leu Asp Thr Val Glu Leu Ala Thr Tyr Thr Val Phe  
 115 120 125  
 Ala Leu His Lys Ser Gly Ser Ser Glu Lys Arg Glu Leu Arg Gln Phe  
 130 135 140  
 Glu Phe Met Ala Trp Pro Asp His Gly Val Pro Glu Tyr Pro Thr Pro  
 145 150 155 160  
 Ile Leu Ala Phe Leu Arg Arg Val Lys Ala Cys Asn Pro Leu Asp Ala  
 165 170 175  
 Gly Pro Met Val Val His Cys Ser Ala Gly Val Gly Arg Thr Gly Cys  
 180 185 190  
 Phe Ile Val Ile Asp Ala Met Leu Glu Arg Met Lys His Glu Lys Thr  
 195 200 205  
 Val Asp Ile Tyr Gly His Val Thr Cys Met Arg Ser Gln Arg Asn Tyr  
 210 215 220  
 Met Val Gln Thr Glu Asp Gln Tyr Val Phe Ile His Glu Ala Leu Leu  
 225 230 235 240  
 Glu Ala Ala Thr Cys Gly His Thr Glu Val Val Ile Asp Ala Met Leu  
 245 250 255  
 Glu Arg Met Lys His Glu Lys Thr Val Asp Ile Tyr Gly His Val Thr  
 260 265 270  
 Cys Met Arg Ser Gln Arg Asn Tyr Met Val Gln Thr Glu Asp Gln Tyr  
 275 280 285  
 Val Phe Ile His Glu Ala Leu Leu Glu Ala Ala Thr Cys Gly His Thr  
 290 295 300  
 Glu Val  
 305

<210> 9

<211> 305

<212> PRT

<213> Homo sapiens

<400> 9

Ser Ala Pro Trp Asp Ser Ala Lys Lys Asp Glu Asn Arg Met Lys Asn  
 1 5 10 15  
 Arg Tyr Gly Asn Ile Ile Ala Tyr Asp His Ser Arg Val Arg Leu Gln  
 20 25 30  
 Thr Ile Glu Gly Asp Thr Asn Ser Asp Tyr Ile Asn Gly Asn Tyr Ile  
 35 40 45  
 Asp Gly Tyr His Arg Pro Asn His Tyr Ile Ala Thr Gln Gly Pro Met  
 50 55 60

Gln Glu Thr Ile Tyr Asp Phe Trp Arg Met Val Trp His Glu Asn Thr  
 65 70 75 80  
 Ala Ser Ile Ile Met Val Thr Asn Leu Val Glu Val Gly Arg Val Lys  
 85 90 95  
 Cys Cys Lys Tyr Trp Pro Asp Asp Thr Glu Ile Tyr Lys Asp Ile Lys  
 100 105 110  
 Val Thr Leu Ile Glu Thr Glu Leu Ala Glu Tyr Val Ile Phe Ala  
 115 120 125  
 Val Glu Lys Arg Gly Val His Glu Ile Arg Glu Ile Arg Gln Phe His  
 130 135 140  
 Phe Thr Gly Trp Pro Asp His Gly Val Pro Tyr His Ala Thr Gly Leu  
 145 150 155 160  
 Leu Gly Phe Val Arg Gln Val Lys Ser Lys Ser Pro Pro Ser Ala Gly  
 165 170 175  
 Pro Leu Val Val His Cys Ser Ala Gly Ala Gly Arg Thr Gly Cys Phe  
 180 185 190  
 Ile Val Ile Asp Ile Met Leu Asp Met Ala Glu Arg Glu Gly Val Val  
 195 200 205  
 Asp Ile Tyr Asn Cys Val Arg Glu Leu Arg Ser Arg Arg Val Asn Met  
 210 215 220  
 Val Gln Thr Glu Glu Gln Tyr Val Phe Ile His Asp Ala Ile Leu Glu  
 225 230 235 240  
 Ala Cys Leu Cys Gly Asp Thr Ser Val Val Ile Asp Ile Met Leu Asp  
 245 250 255  
 Met Ala Glu Arg Glu Gly Val Val Asp Ile Tyr Asn Cys Val Arg Glu  
 260 265 270  
 Leu Arg Ser Arg Arg Val Asn Met Val Gln Thr Glu Glu Gln Tyr Val  
 275 280 285  
 Phe Ile His Asp Ala Ile Leu Glu Ala Cys Leu Cys Gly Asp Thr Ser  
 290 295 300  
 Val  
 305

&lt;10&gt; 10

&lt;11&gt; 310

&lt;12&gt; PET

&lt;13&gt; Homo sapiens

&lt;400&gt; 10

Gln Ala Thr Cys Glu Ala Ala Ser Lys Glu Glu Asn Lys Glu Lys Asn  
 1 5 10 15  
 Arg Tyr Val Asn Ile Leu Pro Tyr Asp His Ser Arg Val His Leu Thr  
 20 25 30  
 Pro Val Glu Gly Val Pro Asp Ser Asp Tyr Ile Asn Ala Ser Phe Ile  
 35 40 45  
 Asn Gly Tyr Gln Glu Lys Asn Lys Phe Ile Ala Ala Gln Gly Pro Lys  
 50 55 60  
 Glu Glu Thr Val Asn Asp Phe Trp Arg Met Ile Trp Glu Gln Asn Thr  
 65 70 75 80  
 Ala Thr Ile Val Met Val Thr Asn Leu Lys Glu Arg Lys Glu Cys Lys  
 85 90 95  
 Cys Ala Gln Tyr Trp Pro Asp Gln Gly Cys Trp Thr Tyr Gly Asn Ile  
 100 105 110  
 Arg Val Ser Val Glu Asp Val Thr Val Leu Val Asp Tyr Thr Val Phe



115	120	125
Cys Ile Gln Gln Val Gly Asp Met Thr Asn Arg Lys Pro Gln Arg Leu		
130	135	140
Ile Thr Gln Phe His Phe Thr Ser Trp Pro Asp Phe Gly Val Pro Phe		
145	150	155
Thr Pro Ile Gly Met Leu Lys Phe Leu Lys Lys Val Lys Ala Cys Asn		
	165	170
Pro Gln Tyr Ala Gly Ala Ile Val Val His Cys Ser Ala Gly Val Gly		
	180	185
Arg Thr Gly Thr Phe Val Val Ile Asp Ala Met Leu Asp Met Met His		
	195	200
Thr Glu Arg Lys Val Asp Val Tyr Gly Phe Val Ser Arg Ile Arg Ala		
	210	215
Gln Arg Cys Gln Met Val Gln Thr Asp Met Gln Tyr Val Phe Ile Tyr		
225	230	235
Gln Ala Leu Leu Glu His Tyr Leu Tyr Gly Asp Thr Glu Leu Val Ile		
	245	250
Asp Ala Met Leu Asp Met Met His Thr Glu Arg Lys Val Asp Val Tyr		
	260	265
Gly Phe Val Ser Arg Ile Arg Ala Gln Arg Cys Gln Met Val Gln Thr		
	275	280
Asp Met Gln Tyr Val Phe Ile Tyr Gln Ala Leu Leu Glu His Tyr Leu		
	290	295
Tyr Gly Asp Thr Glu Leu		
305	310	

&lt;210&gt; 11

&lt;211&gt; 209

&lt;212&gt; FRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 11

Gln Gly Thr Phe Glu Leu Ala Asn Lys Glu Glu Asn Arg Glu Lys Asn		
1	5	10
Arg Tyr Pro Asn Ile Leu Pro Asn Asp His Ser Arg Val Ile Leu Ser		
	20	25
Gln Leu Asp Gly Ile Pro Cys Ser Asp Tyr Ile Asn Ala Ser Tyr Ile		
	35	40
Asp Gly Tyr Lys Glu Lys Asn Lys Phe Ile Ala Ala Gln Gly Pro Lys		
	50	55
Gln Glu Thr Val Asn Asp Phe Trp Arg Met Val Trp Glu Gln Lys Ser		
65	70	75
Ala Thr Ile Val Met Leu Thr Asn Leu Lys Glu Arg Lys Glu Glu Lys		
	85	90
Cys His Gln Tyr Trp Pro Asp Gln Gly Cys Trp Thr Tyr Gly Asn Ile		
	100	105
Arg Val Cys Val Glu Asp Cys Val Val Leu Val Asp Tyr Thr Ile Phe		
	115	120
Cys Ile Gln Pro Gln Leu Pro Asp Gly Cys Lys Ala Pro Arg Leu Val		
	130	135
Ser Gln Leu His Phe Thr Ser Trp Pro Asp Phe Gly Val Pro Phe Thr		
145	150	155
Pro Ile Gly Met Leu Lys Phe Leu Lys Lys Val Lys Thr Leu Asn Pro		
	165	170

Val His Ala Gly Pro Ile Val Val His Cys Ser Ala Gly Val Gly Arg  
 180 185 190  
 Thr Gly Thr Phe Ile Val Ile Asp Ala Met Met Ala Met Met His Ala  
 195 200 205  
 Glu Gln Lys Val Asp Val Phe Glu Phe Val Ser Arg Ile Arg Asn Gln  
 210 215 220  
 Arg Pro Gln Met Val Gln Thr Asp Met Gln Tyr Thr Phe Ile Tyr Gln  
 225 230 235 240  
 Ala Leu Leu Glu Tyr Tyr Leu Tyr Gly Asp Thr Glu Leu Val Ile Asp  
 245 250 255  
 Ala Met Met Ala Met Met His Ala Glu Gln Lys Val Asp Val Phe Glu  
 260 265 270  
 Phe Val Ser Arg Ile Arg Asn Gln Arg Pro Gln Met Val Gln Thr Asp  
 275 280 285  
 Met Gln Tyr Thr Phe Ile Tyr Gln Ala Leu Leu Glu Tyr Tyr Leu Tyr  
 290 295 300  
 Gly Asp Thr Glu Leu  
 305

<210> 12

<211> 309

<212> PRT

<213> Mus musculus

<400> 12

Lys Phe Pro Ile Lys Asp Ala Arg Lys Pro His Asn Gln Asn Lys Asn  
 1 5 10 15  
 Arg Tyr Val Asp Ile Leu Pro Tyr Asp Tyr Asn Arg Val Glu Leu Ser  
 20 25 30  
 Glu Ile Asn Gly Asp Ala Gly Ser Thr Tyr Ile Asn Ala Ser Tyr Ile  
 35 40 45  
 Asp Gly Phe Lys Glu Pro Arg Lys Tyr Ile Ala Ala Gln Gly Pro Arg  
 50 55 60  
 Asp Glu Thr Val Asp Asp Phe Trp Arg Met Ile Trp Glu Gln Lys Ala  
 65 70 75 80  
 Thr Val Ile Val Met Val Thr Arg Cys Glu Glu Gly Asn Arg Asn Lys  
 85 90 95  
 Cys Ala Glu Tyr Trp Pro Ser Met Glu Glu Gly Thr Arg Ala Phe Lys  
 100 105 110  
 Asp Ile Val Val Thr Ile Asn Asp His Lys Arg Cys Pro Asp Tyr Ile  
 115 120 125  
 Ile Leu Asn Val Ala His Lys Lys Glu Lys Ala Thr Gly Arg Glu Val  
 130 135 140  
 Thr His Ile Gln Phe Thr Ser Trp Pro Asp His Gly Val Pro Glu Asp  
 145 150 155 160  
 Pro His Leu Leu Leu Lys Leu Arg Arg Arg Val Asn Ala Phe Ser Asn  
 165 170 175  
 Phe Phe Ser Gly Pro Ile Val Val His Cys Ser Ala Gly Val Gly Arg  
 180 185 190  
 Thr Gly Thr Tyr Ile Gly Ile Asp Ala Met Leu Glu Gly Leu Glu Ala  
 195 200 205  
 Glu Gly Lys Val Asp Val Tyr Gly Tyr Val Val Lys Leu Arg Arg Gln  
 210 215 220  
 Arg Cys Leu Met Val Gln Val Glu Ala Gln Tyr Ile Leu Ile His Gln

245                      240                      235                      240  
 Ala Leu Val Glu Tyr Asn Gln Phe Gly Glu Thr Glu Val Gly Ile Asp  
                                  245                      250                      255  
 Ala Met Leu Glu Gly Leu Glu Ala Glu Gly Lys Val Asp Val Tyr Gly  
                                  260                      265                      270  
 Tyr Val Val Lys Leu Arg Arg Gln Arg Cys Leu Met Val Gln Val Glu  
                                  275                      280                      285  
 Ala Gln Tyr Ile Leu Ile His Gln Ala Leu Val Glu Tyr Asn Gln Phe  
                                  290                      295                      300  
 Gly Glu Thr Glu Val  
 305

<210> 13  
 <211> 325  
 <212> PRT  
 <213> Homo sapiens

<100> 13  
 Leu Tyr Ser Arg Lys Glu Gly Gln Arg Gln Gln Asn Lys Asn Lys Asn  
   1                      5                      10                      15  
 Arg Tyr Lys Asn Ile Leu Pro Phe Asp His Thr Arg Val Val Leu His  
                                  20                      25                      30  
 Asp Gly Asp Pro Asn Glu Pro Val Ser Asp Tyr Ile Asn Ala Asn Ile  
                                  35                      40                      45  
 Ile Met Pro Glu Phe Glu Thr Lys Cys Asn Asn Ser Lys Pro Lys Lys  
                                  50                      55                      60  
 Ser Tyr Ile Ala Thr Gln Gly Cys Leu Gln Asn Thr Val Asn Asp Phe  
                                  65                      70                      75                      80  
 Trp Arg Met Val Phe Gln Glu Asn Ser Arg Val Ile Val Met Thr Thr  
                                  85                      90                      95  
 Lys Glu Val Glu Arg Gly Lys Ser Lys Cys Val Lys Tyr Trp Pro Asp  
                                  100                      105                      110  
 Glu Tyr Ala Leu Lys Glu Tyr Gly Val Met Arg Val Arg Asn Val Lys  
                                  115                      120                      125  
 Glu Ser Ala Ala His Asp Tyr Thr Leu Leu Lys Leu Ser Lys Val Gly  
                                  130                      135                      140  
 Gln Gly Asn Thr Glu Arg Thr Val Trp Gln Tyr His Phe Arg Thr Trp  
                                  145                      150                      155                      160  
 Pro Asp His Gly Val Pro Ser Asp Pro Gly Gly Val Leu Asp Phe Leu  
                                  165                      170                      175  
 Glu Glu Val His His Lys Gln Glu Ser Ile Met Asp Ala Gly Pro Val  
                                  180                      185                      190  
 Val Val His Cys Ser Ala Gly Ile Gly Arg Thr Gly Thr Phe Ile Val  
                                  195                      200                      205  
 Ile Asp Ile Leu Ile Asp Ile Ile Arg Glu Lys Gly Val Asp Cys Asp  
                                  210                      215                      220  
 Ile Asp Val Pro Lys Thr Ile Gln Met Val Arg Ser Gln Arg Ser Gly  
                                  225                      230                      235                      240  
 Met Val Gln Thr Glu Ala Gln Tyr Arg Phe Ile Tyr Met Ala Val Gln  
                                  245                      250                      255  
 His Tyr Ile Glu Thr Leu Gln Arg Arg Ile Val Ile Asp Ile Leu Ile  
                                  260                      265                      270  
 Asp Ile Ile Arg Glu Lys Gly Val Asp Cys Asp Ile Asp Val Pro Lys  
                                  275                      280                      285

Thr Ile Gln Met Val Arg Ser Gln Arg Ser Gly Met Val Gln Thr Glu  
 290 295 300  
 Ala Gln Tyr Arg Phe Ile Tyr Met Ala Val Gln His Tyr Ile Glu Thr  
 305 310 315 320  
 Leu Gln Arg Arg Ile  
 325

<210> 14

<211> 322

<212> PRT

<213> Homo sapiens

<400> 14

Leu His Gln Arg Leu Glu Gly Gln Arg Pro Glu Asn Lys Gly Lys Asn  
 1 5 10 15  
 Arg Tyr Lys Asn Ile Leu Pro Phe Asp His Ser Arg Val Ile Leu Gln  
 20 25 30  
 Gly Arg Asp Ser Asn Ile Pro Gly Ser Asp Tyr Ile Asn Ala Asn Tyr  
 35 40 45  
 Ile Lys Asn Gln Leu Leu Gly Pro Asp Glu Asn Ala Lys Thr Tyr Ile  
 50 55 60  
 Ala Ser Gln Gly Cys Leu Glu Ala Thr Val Asn Asp Phe Trp Gln Met  
 65 70 75 80  
 Ala Trp Gln Glu Asn Ser Arg Val Ile Val Met Thr Thr Arg Glu Val  
 85 90 95  
 Glu Lys Gly Arg Asn Lys Cys Val Pro Tyr Trp Pro Glu Val Gly Met  
 100 105 110  
 Gln Arg Ala Tyr Gly Pro Tyr Ser Val Thr Asn Cys Gly Glu His Asp  
 115 120 125  
 Thr Thr Gln Tyr Lys Leu Leu Gln Val Ser Pro Leu Asp Asn Gly Asp  
 130 135 140  
 Leu Ile Arg Glu Ile Trp His Tyr Gln Tyr Leu Ser Trp Pro Asp His  
 145 150 155 160  
 Gly Val Pro Ser Glu Pro Gly Gly Val Leu Ser Phe Leu Asp Gln Ile  
 165 170 175  
 Asn Gln Arg Gln Glu Ser Leu Pro His Ala Gly Pro Ile Ile Val His  
 180 185 190  
 Cys Ser Ala Gly Ile Gly Arg Thr Gly Thr Ile Ile Val Ile Asp Met  
 195 200 205  
 Leu Met Glu Asn Ile Ser Thr Lys Gly Leu Asp Cys Asp Ile Asp Ile  
 210 215 220  
 Gln Lys Thr Ile Gln Met Val Arg Ala Gln Arg Ser Gly Met Val Gln  
 225 230 235 240  
 Thr Glu Ala Gln Tyr Lys Phe Ile Tyr Val Ala Ile Ala Gln Phe Ile  
 245 250 255  
 Glu Thr Thr Lys Lys Lys Leu Val Ile Asp Met Leu Met Glu Asn Ile  
 260 265 270  
 Ser Thr Lys Gly Leu Asp Cys Asp Ile Asp Ile Gln Lys Thr Ile Gln  
 275 280 285  
 Met Val Arg Ala Gln Arg Ser Gly Met Val Gln Thr Glu Ala Gln Tyr  
 290 295 300  
 Lys Phe Ile Tyr Val Ala Ile Ala Gln Phe Ile Glu Thr Thr Lys Lys  
 305 310 315 320  
 Lys Leu

<210> 15  
 <211> 319  
 <212> PPT  
 <213> Homo sapiens

<400> 15

Asn	Gln	Ser	Cys	Asp	Ile	Ala	Leu	Leu	Pro	Gln	Asn	Arg	Gly	Lys	Asn
1			5						10					15	
Arg	Tyr	Asn	Asn	Ile	Leu	Pro	Tyr	Asp	Ala	Thr	Arg	Val	Lys	Leu	Ser
		20						25					30		
Asn	Val	Asp	Asp	Asp	Pro	Cys	Ser	Asp	Tyr	Ile	Asn	Ala	Ser	Tyr	Ile
	35						40					45			
Pro	Gly	Asn	Asn	Phe	Arg	Arg	Glu	Tyr	Ile	Val	Thr	Gln	Gly	Pro	Leu
	50					55					60				
Pro	Gly	Thr	Lys	Asp	Asp	Phe	Trp	Lys	Met	Val	Trp	Gln	Gln	Asn	Val
65				70				75						80	
His	Asn	Ile	Val	Met	Val	Thr	Gln	Cys	Val	Glu	Lys	Gly	Arg	Val	Lys
			85					90						95	
Cys	Asp	His	Tyr	Trp	Pro	Ala	Asp	Gln	Asp	Ser	Leu	Tyr	Tyr	Gly	Asp
	100						105							110	
Leu	Ile	Leu	Gln	Met	Leu	Ser	Glu	Ser	Val	Leu	Pro	Glu	Trp	Thr	Ile
	115						120					125			
Phe	Lys	Ile	Cys	Gly	Glu	Gln	Gln	Leu	Asp	Ala	His	Arg	Leu	Ile	Arg
	130						135					140			
His	Phe	His	Tyr	Thr	Val	Trp	Pro	Asp	His	Gly	Val	Pro	Glu	Thr	Thr
	145					150				155					160
Gln	Ser	Leu	Ile	Gln	Phe	Val	Arg	Thr	Val	Arg	Asp	Tyr	Ile	Asn	Arg
			165					170						175	
Ser	Pro	Gly	Ala	Gly	Pro	Thr	Val	Val	His	Cys	Ser	Ala	Gly	Val	Gly
	180							185						190	
Arg	Thr	Gly	Thr	Phe	Ile	Ala	Leu	Asp	Arg	Ile	Leu	Gln	Gln	Leu	Asp
	195						200						205		
Ser	Lys	Asp	Ser	Val	Asp	Ile	Tyr	Gly	Ala	Val	His	Asp	Leu	Arg	Leu
	210					215						220			
His	Arg	Val	His	Met	Val	Gln	Thr	Glu	Cys	Gln	Tyr	Val	Tyr	Leu	His
225					230					235					240
Gln	Cys	Val	Arg	Asp	Val	Leu	Arg	Ala	Arg	Lys	Leu	Arg	Ser	Ala	Leu
			245						250					255	
Asp	Arg	Ile	Leu	Gln	Gln	Leu	Asp	Ser	Lys	Asp	Ser	Val	Asp	Ile	Tyr
			260					265					270		
Gly	Ala	Val	His	Asp	Leu	Arg	Leu	His	Arg	Val	His	Met	Val	Gln	Thr
	275						280					285			
Glu	Cys	Gln	Tyr	Val	Tyr	Leu	His	Gln	Cys	Val	Arg	Asp	Val	Leu	Arg
	290					295						300			
Ala	Arg	Lys	Leu	Arg	Ser										
305					310										

<210> 16  
 <211> 399  
 <212> PRT  
 <213> *Prosephila melanogaster*

&lt;400&gt; 16

Asp Gln Pro Cys Thr Phe Ala Asp Leu Pro Cys Asn Arg Pro Lys Asn  
 1 5 10 15  
 Arg Phe Thr Asn Ile Leu Pro Tyr Asp His Ser Arg Phe Lys Leu Gln  
 20 25 30  
 Pro Val Asp Asp Asp Glu Gly Ser Asp Tyr Ile Asn Ala Asn Tyr Val  
 35 40 45  
 Pro Gly His Asn Ser Pro Arg Glu Phe Ile Val Thr Gln Gly Pro Leu  
 50 55 60  
 His Ser Thr Arg Asp Asp Phe Trp Arg Met Cys Trp Glu Ser Asn Ser  
 65 70 75 80  
 Arg Ala Ile Val Met Leu Thr Arg Cys Phe Glu Lys Gly Arg Glu Lys  
 85 90 95  
 Cys Asp Gln Tyr Trp Pro Asn Asp Thr Val Pro Val Phe Tyr Gly Asp  
 100 105 110  
 Ile Lys Val Gln Ile Leu Asn Asp Ser His Tyr Ala Asp Trp Val Met  
 115 120 125  
 Phe Met Leu Cys Arg Gly Ser Glu Gln Arg Ile Leu Arg His Phe His  
 130 135 140  
 Phe Thr Thr Trp Pro Asp Phe Gly Val Pro Asn Pro Pro Gln Thr Leu  
 145 150 155 160  
 Val Arg Phe Val Arg Ala Phe Arg Asp Arg Ile Cys Ala Glu Gln Arg  
 165 170 175  
 Pro Ile Val Val His Cys Ser Ala Gly Val Gly Arg Ser Gly Thr Phe  
 180 185 190  
 Ile Thr Leu Asp Arg Ile Leu Gln Gln Ile Asn Thr Ser Asp Tyr Val  
 195 200 205  
 Asp Ile Phe Gly Ile Val Tyr Ala Met Arg Lys Glu Arg Val Trp Met  
 210 215 220  
 Val Gln Thr Glu Gln Gln Tyr Ile Cys Ile His Gln Cys Leu Leu Ala  
 225 230 235 240  
 Val Leu Glu Gly Lys Glu Asn Ile Val Gly Pro Thr Leu Asp Arg Ile  
 245 250 255  
 Leu Gln Gln Ile Asn Thr Ser Asp Tyr Val Asp Ile Phe Gly Ile Val  
 260 265 270  
 Tyr Ala Met Arg Glu Lys Arg Val Trp Met Val Gln Thr Glu Gln Gln  
 275 280 285  
 Tyr Ile Cys Ile His Gln Cys Leu Leu Ala Val Leu Glu Gly Lys Glu  
 290 295 300  
 Asn Ile Val Gly Pro  
 305

&lt;210&gt; 17

&lt;211&gt; 313

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 17

Ser Gln Ser Gln Met Val Ala Ser Ala Ser Glu Asn Asn Ala Lys Asn  
 1 5 10 15  
 Arg Tyr Arg Asn Val Leu Pro Tyr Asp Trp Ser Arg Val Phe Leu Lys  
 20 25 30  
 Pro Ile His Glu Glu Pro Gly Ser Asp Tyr Ile Asn Ala Ser Ile Met  
 35 40 45

Pro Gly Leu Trp Ser Pro Gln Glu Phe Ile Ala Thr Gln Gly Trp Leu  
 50 55 60  
 Pro Gln Thr Val Gly Asp Phe Trp Arg Leu Val Trp Glu Gln Gln Ser  
 65 70 75 80  
 His Thr Leu Val Met Leu Thr Asn Cys Met Glu Ala Gly Arg Val Lys  
 85 90 95  
 Cys Glu His Tyr Trp Pro Leu Asp Ser Gln Pro Cys Thr His Gly His  
 100 105 110  
 Leu Arg Val Thr Leu Val Gly Glu Glu Val Met Glu Asn Trp Thr Val  
 115 120 125  
 Leu Leu Leu Leu Gln Val Glu Glu Gln Lys Thr Leu Ser Val Arg Gln  
 130 135 140  
 Phe His Tyr Gln Ala Trp Pro Asp His Gly Val Pro Ser Ser Pro Asp  
 145 150 155 160  
 Thr Leu Leu Ala Phe Trp Arg Met Leu Arg Gln Trp Leu Asp Gln Thr  
 165 170 175  
 Met Glu Gly Gly Pro Pro Ile Val His Cys Ser Ala Gly Val Gly Arg  
 180 185 190  
 Thr Gly Thr Leu Ile Ala Leu Asp Val Leu Leu Arg Gln Leu Gln Ser  
 195 200 205  
 Glu Gly Leu Leu Gly Pro Phe Ser Phe Val Arg Lys Met Arg Glu Ser  
 210 215 220  
 Arg Pro Leu Met Val Gln Thr Glu Ala Gln Tyr Val Phe Leu His Gln  
 225 230 235 240  
 Cys Ile Cys Gly Ser Ser Asn Ser Gln Pro Arg Pro Gln Pro Arg Ala  
 245 250 255  
 Leu Asp Val Leu Leu Arg Gln Leu Gln Ser Glu Gly Leu Leu Gly Pro  
 260 265 270  
 Phe Ser Phe Val Arg Lys Met Arg Glu Ser Arg Pro Leu Met Val Gln  
 275 280 285  
 Thr Glu Ala Gln Tyr Val Phe Leu His Gln Cys Ile Cys Gly Ser Ser  
 290 295 300  
 Asn Ser Gln Pro Arg Pro Gln Pro Arg  
 305 310

&lt;.10&gt; 18

&lt;.11&gt; 291

&lt;.12&gt; PRT

&lt;.13&gt; Rattus norvegicus

&lt;400&gt; 18

Phe Val Asp Pro Lys Glu Tyr Asp Ile Pro Gly Leu Val Arg Lys Asn  
 1 5 10 15  
 Arg Tyr Lys Thr Ile Leu Pro Asn Pro His Ser Arg Val Arg Leu Thr  
 20 25 30  
 Ser Pro Asp Pro Glu Asp Pro Leu Ser Ser Tyr Ile Asn Ala Asn Tyr  
 35 40 45  
 Ile Arg Gly Tyr Asn Gly Glu Lys Val Tyr Ile Ala Thr Gln Gly  
 50 55 60  
 Pro Ile Val Ser Thr Val Val Asp Phe Trp Arg Met Val Trp Glu Glu  
 65 70 75 80  
 Arg Thr Pro Ile Ile Val Met Ile Thr Asn Ile Glu Glu Met Asn Glu  
 85 90 95  
 Lys Cys Thr Glu Tyr Trp Pro Glu Glu Gln Val Val His Asp Gly Val

	105		105		110										
Glu	Ile	Thr	Val	Gln	Lys	Val	Ile	His	Thr	Glu	Asp	Tyr	Arg	Leu	Ile
	115						120						125		
Ser	Leu	Arg	Arg	Gly	Thr	Glu	Glu	Arg	Gly	Leu	Lys	His	Tyr	Trp	Phe
	130						135						140		
Thr	Ser	Trp	Pro	Asp	Gln	Lys	Thr	Pro	Asp	Arg	Ala	Pro	Pro	Leu	Leu
	145					150				155				160	
His	Leu	Val	Arg	Glu	Val	Glu	Glu	Ala	Ala	Gln	Gln	Glu	Gly	Pro	His
				165					170					175	
Cys	Ser	Pro	Ile	Ile	Val	His	Cys	Ser	Ala	Gly	Ile	Gly	Arg	Thr	Gly
			180				185						190		
Cys	Phe	Ile	Ala	Thr	Ser	Ile	Cys	Cys	Gln	Gln	Leu	Arg	Arg	Glu	Gly
	195						200					205			
Val	Val	Asp	Ile	Leu	Lys	Thr	Thr	Cys	Gln	Leu	Arg	Gln	Asp	Arg	Gly
	210					215					220				
Gly	Met	Ile	Gln	Thr	Cys	Glu	Gln	Tyr	Gln	Phe	Val	His	His	Ala	Met
	225				230				235						240
Ser	Leu	Tyr	Ala	Thr	Ser	Ile	Cys	Cys	Gln	Gln	Leu	Arg	Arg	Glu	Gly
			245					250						255	
Val	Val	Asp	Ile	Leu	Lys	Thr	Thr	Cys	Gln	Leu	Arg	Gln	Asp	Arg	Gly
		260					265					270			
Gly	Met	Ile	Gln	Thr	Cys	Glu	Gln	Tyr	Gln	Phe	Val	His	His	Ala	Met
	275					280						285			
Ser	Leu	Tyr													
	290														

&lt;210&gt; 19

&lt;211&gt; 313

&lt;212&gt; PRT

<213> *Drosophila melanogaster*

&lt;400&gt; 19

Asp	Arg	Thr	Thr	Lys	Asn	Ser	Asp	Leu	Lys	Glu	Asn	Ala	Cys	Lys	Asn
1				5					10					15	
Arg	Tyr	Pro	Asp	Ile	Lys	Ala	Tyr	Asp	Gln	Thr	Arg	Val	Lys	Leu	Ala
		20						25					30		
Val	Ile	Asn	Gly	Leu	Gln	Thr	Thr	Asp	Tyr	Ile	Asn	Ala	Asn	Phe	Val
	35					40					45				
Ile	Gly	Tyr	Lys	Glu	Arg	Lys	Lys	Phe	Ile	Cys	Ala	Gln	Gly	Pro	Met
	50					55				60					
Glu	Ser	Thr	Ile	Asp	Asp	Phe	Trp	Arg	Met	Ile	Trp	Glu	Gln	His	Leu
	65			70					75					80	
Glu	Ile	Ile	Val	Ile	Leu	Thr	Asn	Leu	Glu	Glu	Tyr	Asn	Lys	Ala	Lys
			85						90				95		
Cys	Ala	Lys	Tyr	Trp	Pro	Glu	Lys	Val	Phe	Asp	Thr	Lys	Gln	Phe	Gly
	100						105						110		
Asp	Ile	Leu	Val	Lys	Phe	Ala	Gln	Glu	Arg	Lys	Thr	Gly	Asp	Tyr	Ile
	115						120					125			
Glu	Leu	Asn	Val	Ser	Lys	Asn	Lys	Ala	Asn	Val	Gly	Glu	Glu	Glu	Asp
	130					135					140				
Arg	Arg	Gln	Ile	Thr	Gln	Tyr	His	Tyr	Leu	Thr	Trp	Lys	Asp	Phe	Met
	145				150				155					160	
Ala	Pro	Glu	His	Pro	His	Gly	Ile	Ile	Lys	Phe	Ile	Arg	Gln	Ile	Asn
			165					170						175	



Ser Val Tyr Ser Leu Gln Arg Gly Pro Ile Leu Val His Cys Ser Ala  
 180 185 190  
 Gly Val Gly Arg Thr Gly Thr Leu Val Ala Leu Asp Ser Leu Ile Gln  
 195 200 205  
 Gln Leu Glu Glu Glu Asp Ser Val Ser Ile Tyr Asn Thr Val Cys Asp  
 210 215 220  
 Leu Arg His Gln Arg Asn Phe Leu Val Gln Ser Leu Lys Gln Tyr Ile  
 225 230 235 240  
 Phe Leu Tyr Arg Ala Leu Leu Asp Thr Gly Thr Phe Gly Asn Thr Asp  
 245 250 255  
 Ile Ala Leu Asp Ser Leu Ile Gln Gln Leu Glu Glu Glu Asp Ser Val  
 260 265 270  
 Ser Ile Tyr Asn Thr Val Cys Asp Leu Arg His Gln Arg Asn Phe Leu  
 275 280 285  
 Val Gln Ser Leu Lys Gln Tyr Ile Phe Leu Tyr Arg Ala Leu Leu Asp  
 290 295 300  
 Thr Gly Thr Phe Gly Asn Thr Asp Ile  
 305 310

<210> 20

<211> 307

<212> PRT

<213> Homo sapiens

<400> 20

Val Gly Thr Phe His Cys Ser Met Ser Pro Gly Asn Leu Glu Lys Asn  
 1 5 10 15  
 Arg Tyr Gly Asp Val Pro Cys Leu Asp Gln Thr Arg Val Lys Leu Thr  
 20 25 30  
 Lys Arg Ser Gly His Thr Gln Thr Asp Tyr Ile Asn Ala Ser Phe Met  
 35 40 45  
 Asp Gly Tyr Lys Gln Lys Asn Ala Tyr Ile Gly Thr Gln Gly Pro Leu  
 50 55 60  
 Glu Asn Thr Tyr Arg Asp Phe Trp Leu Met Val Trp Glu Gln Lys Val  
 65 70 75 80  
 Leu Val Ile Val Met Thr Thr Arg Phe Glu Glu Gly Gly Arg Arg Lys  
 85 90 95  
 Cys Gly Gln Tyr Trp Pro Leu Glu Lys Asp Ser Arg Ile Arg Phe Gly  
 100 105 110  
 Phe Leu Thr Val Thr Asn Leu Gly Val Glu Asn Met Asn His Tyr Lys  
 115 120 125  
 Lys Leu Glu Ile His Asn Thr Glu Glu Arg Gln Lys Arg Gln Val Thr  
 130 135 140  
 His Phe Gln Phe Leu Ser Trp Pro Asp Tyr Gly Val Pro Ser Ser Ala  
 145 150 155 160  
 Ala Ser Leu Ile Asp Phe Leu Arg Val Val Arg Asn Gln Gln Ser Leu  
 165 170 175  
 Ala Val Ser Asn Met Gly Ala Arg Ser Lys Gly Gln Cys Pro Glu Pro  
 180 185 190  
 Pro Ile Val Val His Cys Ser Ala Gly Ile Gly Arg Thr Gly Thr Phe  
 195 200 205  
 Cys Ser Leu Asp Ile Cys Leu Ala Gln Leu Glu Glu Leu Gly Thr Leu  
 210 215 220  
 Asn Val Phe Gln Thr Val Ser Arg Met Arg Thr Gln Arg Ala Phe Ser

225 230 235 240  
 Ile Gln Thr Pro Glu Gln Tyr Tyr Phe Cys Tyr Lys Ala Ile Leu Glu  
 245 250 255  
 Phe Ala Ser Leu Asp Ile Cys Leu Ala Gln Leu Glu Glu Leu Gly Thr  
 260 265 270  
 Leu Asn Val Phe Gln Thr Val Ser Arg Met Arg Thr Gln Arg Ala Phe  
 275 280 285  
 Ser Ile Gln Thr Pro Glu Gln Tyr Tyr Phe Cys Tyr Lys Ala Ile Leu  
 290 295 300  
 Glu Phe Ala  
 305

<210> 21

<211> 312

<212> PRT

<213> Homo sapiens

<400> 21

1 5 10 15  
 Ile Tyr Pro Thr Ala Thr Gly Glu Lys Glu Glu Asn Val Lys Lys Asn  
 Arg Tyr Lys Asp Ile Leu Pro Phe Asp His Ser Arg Val Lys Leu Thr  
 20 25 30  
 Leu Lys Thr Pro Ser Gln Asp Ser Asp Tyr Ile Asn Ala Asn Phe Ile  
 35 40 45  
 Lys Gly Val Tyr Gly Pro Lys Ala Tyr Val Ala Thr Gln Gly Pro Leu  
 50 55 60  
 Ala Asn Thr Val Ile Asp Phe Trp Arg Met Val Trp Glu Tyr Asn Val  
 65 70 75 80  
 Val Ile Ile Val Met Ala Cys Arg Glu Phe Glu Met Gly Arg Lys Lys  
 85 90 95  
 Cys Glu Arg Tyr Trp Pro Leu Tyr Gly Glu Asp Pro Ile Thr Phe Ala  
 100 105 110  
 Pro Phe Lys Ile Ser Cys Glu Asp Glu Gln Ala Arg Thr Asp Tyr Phe  
 115 120 125  
 Ile Leu Leu Leu Glu Phe Gln Asn Glu Ser Arg Arg Leu Tyr Gln Phe  
 130 135 140  
 His Tyr Val Asn Trp Pro Asp His Asp Val Pro Ser Ser Phe Asp Ser  
 145 150 155 160  
 Ile Leu Asp Met Ile Ser Leu Met Arg Lys Tyr Gln Glu His Glu Asp  
 165 170 175  
 Val Pro Ile Cys Ile His Cys Ser Ala Gly Cys Gly Arg Thr Gly Ala  
 180 185 190  
 Ile Cys Ala Ile Asp Tyr Thr Trp Asn Leu Leu Lys Ala Gly Lys Ile  
 195 200 205  
 Pro Glu Glu Phe Asn Val Phe Asn Leu Ile Gln Glu Met Arg Thr Gln  
 210 215 220  
 Arg His Ser Ala Val Gln Thr Lys Glu Gln Tyr Glu Leu Val His Arg  
 225 230 235 240  
 Ala Ile Ala Gln Leu Phe Glu Lys Gln Leu Gln Leu Tyr Ala Ile Asp  
 245 250 255  
 Tyr Thr Trp Asn Leu Leu Lys Ala Gly Lys Ile Pro Glu Glu Phe Asn  
 260 265 270  
 Val Phe Asn Leu Ile Gln Glu Met Arg Thr Gln Arg His Ser Ala Val  
 275 280 285

Gln Thr Lys Gln Gln Tyr Gln Leu Val His Arg Ala Ile Ala Gln Leu  
 290 295 300  
 Phe Gln Lys Gln Leu Gln Leu Tyr  
 305 310

<210> 22

<211> 291

<212> PRT

<213> Homo sapiens

<400> 22

Gly Leu Ala Ile Thr Phe Ala Lys Leu Pro Gln Asn Leu Asp Lys Asn  
 1 5 10 15  
 Arg Tyr Lys Asp Val Leu Pro Tyr Asp Thr Thr Arg Val Leu Leu Gln  
 20 25 30  
 Gly Asn Glu Asp Tyr Ile Asn Ala Ser Tyr Val Asn Met Glu Ile Pro  
 35 40 45  
 Ala Ala Asn Ile Val Asn Lys Tyr Ile Ala Thr Gln Gly Pro Leu Pro  
 50 55 60  
 His Thr Cys Ala Gln Ile Thr Gln Val Val Trp Asp Gln Lys Leu Ser  
 65 70 75 80  
 Leu Ile Val Met Leu Thr Thr Leu Thr Glu Arg Gly Arg Thr Lys Cys  
 85 90 95  
 His Gln Tyr Trp Pro Asp Pro Pro Asp Val Met Asn His Gly Gly Phe  
 100 105 110  
 His Ile Gln Cys Gln Ser Glu Asp Cys Thr Ile Ala Tyr Val Ser Met  
 115 120 125  
 Leu Val Thr Asn Thr Gln Thr Gly Glu Glu His Thr Val Thr His Leu  
 130 135 140  
 Gln Tyr Val Ala Trp Pro Asp His Gly Ile Pro Asp Asp Ser Ser Asp  
 145 150 155 160  
 Phe Leu Glu Phe Val Asn Tyr Val Arg Ser Leu Arg Val Asp Ser Glu  
 165 170 175  
 Pro Val Leu Val His Cys Ser Ala Gly Ile Gly Arg Thr Gly Val Leu  
 180 185 190  
 Val Thr Met Glu Thr Ala Met Cys Leu Thr Glu Arg Asn Leu Pro Ile  
 195 200 205  
 Tyr Pro Leu Asp Ile Val Arg Lys Met Arg Asp Gln Arg Ala Met Met  
 210 215 220  
 Val Gln Thr Ser Ser Gln Tyr Lys Phe Val Cys Glu Ala Ile Leu Arg  
 225 230 235 240  
 Val Tyr Thr Met Glu Thr Ala Met Cys Leu Thr Glu Arg Asn Leu Pro  
 245 250 255  
 Ile Tyr His Leu Asp Ile Val Arg Lys Met Arg Asp Gln Arg Ala Met  
 260 265 270  
 Met Val Gln Thr Ser Ser Gln Tyr Lys Phe Val Cys Glu Ala Ile Leu  
 275 280 285  
 Arg Val Tyr  
 290

<210> 23

<211> 311

<212> PRT

<213> Canis familiaris familiaris

<400> 23

Pro Ser Glu Thr Ser Glu Gly Asp Lys Lys His Asn Thr Ser Lys Asn  
1 5 10 15  
Arg Tyr Thr Asn Ile Leu Pro Val Asn His Thr Arg Val Gln Leu Lys  
20 25 30  
Lys Ile Gln Asp Lys Glu Gly Ser Asp Tyr Ile Asn Ala Asn Tyr Ile  
35 40 45  
Asp Gly Ala Tyr Pro Lys Gln Phe Ile Cys Thr Gln Gly Pro Leu Pro  
50 55 60  
Asn Thr Ile Ala Asp Phe Trp Arg Met Val Trp Glu Asn Arg Cys Arg  
65 70 75 80  
Ile Ile Val Met Leu Ser Arg Glu Ser Glu Gly Ser Glu Asn Cys Arg  
85 90 95  
Ile Lys Cys Asp Arg Tyr Trp Pro Gln Gln Ile Gly Gly Glu Gln Phe  
100 105 110  
Ser Ile Tyr Gly Asn Gly Asn Glu Val Phe Gly Thr Tyr Ser Val Glu  
115 120 125  
Leu Val Glu Val Ile Gln Cys Arg Glu His Ile Thr Arg Asn Ile Arg  
130 135 140  
Leu Thr Phe Glu Gly Glu Thr Arg Asp Ile Thr Gln Tyr Gln Tyr Glu  
145 150 155 160  
Gly Trp Pro Asp His Asn Ile Pro Asp His Thr Gln Pro Phe Arg Gln  
165 170 175  
Leu Leu His Ser Ile Thr Asn Arg Gln Asn Gln Ile Ile Pro Ser Ser  
180 185 190  
Asp Arg Asn Val Pro Ile Ile Val His Cys Ser Ala Gly Val Gly Arg  
195 200 205  
Thr Gly Thr Phe Cys Thr Ala Val Ile Met Met Lys Lys Leu Asp His  
210 215 220  
Tyr Phe Lys Gln Leu Asp Tyr Asn Ser Arg Ile Asp Phe Asn Leu Phe  
225 230 235 240  
Ser Ile Val Leu Lys Leu Arg Glu Gln Arg Pro Gly Met Val Gln Gln  
245 250 255  
Leu Glu Gln Tyr Leu Phe Cys Tyr Lys Thr Ile Leu Asp Glu Ile Tyr  
260 265 270  
His Arg Leu Asn Cys Thr Ala Val Ile Met Met Lys Lys Leu Asp His  
275 280 285  
Tyr Phe Lys Gln Leu Asp Tyr Asn Ser Arg Ile Asp Phe Asn Leu Phe  
290 295 300  
Ser Ile Val Leu Lys Leu Arg Glu Gln Arg Pro Gly Met Val Gln Gln  
305 310 315 320  
Leu Glu Gln Tyr Leu Phe Cys Tyr Lys Thr Ile Leu Asp Glu Ile Tyr  
325 330 335  
His Arg Leu Asn Cys  
340

<210> 24

<211> 312

<212> PRT

<213> Schistosoma japonicum protein

<400> 24

Ser Trp Ser Thr Val Arg Ser Leu Ser Asn Thr Ser Tyr Lys Lys Asn

1 5 10 15  
 Arg Tyr Thr Asp Ile Val Pro Tyr Asn Cys Thr Arg Val His Leu Lys  
 20 25 30  
 Arg Thr Ser Pro Ser Glu Leu Asp Tyr Ile Asn Ala Ser Phe Ile Lys  
 35 40 45  
 Thr Glu Thr Ser Asn Tyr Ile Ala Cys Gln Gly Ser Ile Ser Arg Ser  
 50 55 60  
 Ile Ser Asp Phe Trp His Met Val Trp Asp Asn Val Glu Asn Ile Gly  
 65 70 75 80  
 Thr Ile Val Met Leu Gly Ser Leu Phe Glu Ala Gly Arg Glu Met Cys  
 85 90 95  
 Thr Ala Tyr Trp Pro Ser Asn Gly Ile Gly Asp Lys Gln Val Tyr Gly  
 100 105 110  
 Asp Tyr Cys Val Lys Gln Ile Ser Glu Glu Asn Val Asp Asn Ser Arg  
 115 120 125  
 Phe Ile Leu Phe Glu Ile Gln Asn Ala Asn Phe Pro Ser Val Lys Lys  
 130 135 140  
 Val His His Tyr Gln Tyr Pro Asn Trp Ser Asp Cys Asn Ser Pro Glu  
 145 150 155  
 Asn Val Lys Ser Met Val Glu Phe Leu Lys Tyr Val Asn Asn Ser His  
 165 170 175  
 Gly Ser Gly Asn Thr Ile Val His Cys Ser Ala Gly Val Gly Arg Thr  
 180 185 190  
 Gly Thr Phe Ile Val Leu Asp Thr Ile Leu Arg Phe Pro Glu Ser Lys  
 195 200 205  
 Leu Ser Gly Phe Asn Pro Ser Val Ala Asp Ser Ser Asp Val Val Phe  
 210 215 220  
 Gln Leu Val Asp His Ile Arg Lys Gln Arg Met Lys Met Val Gln Thr  
 225 230 235 240  
 Phe Thr Gln Phe Lys Tyr Val Tyr Asp Leu Ile Asp Ser Leu Val Leu  
 245 250 255  
 Asp Thr Ile Leu Arg Phe Pro Glu Ser Lys Leu Ser Gly Phe Asn Pro  
 260 265 270  
 Ser Val Ala Asp Ser Ser Asp Val Val Phe Gln Leu Val Asp His Ile  
 275 280 285  
 Arg Lys Gln Arg Met Lys Met Val Gln Thr Phe Thr Gln Phe Lys Tyr  
 290 295 300  
 Val Tyr Asp Leu Ile Asp Ser Leu  
 305 310

&lt;210&gt; 25

&lt;211&gt; 307

&lt;212&gt; PRT

&lt;213&gt; Schizosaccharomyces pombe

&lt;400&gt; 25

Trp Cys Cys Leu Ala Ser Ser Arg Ser Thr Ser Ile Ser Arg Lys Asn  
 1 5 10 15  
 Arg Tyr Thr Asp Ile Val Pro Tyr Asp Lys Thr Arg Val Arg Leu Ala  
 20 25 30  
 Val Pro Lys Gly Cys Ser Asp Tyr Ile Asn Ala Ser His Ile Asp Val  
 35 40 45  
 Gly Asn Lys Lys Tyr Ile Ala Cys Gln Ala Pro Lys Pro Gly Thr Leu  
 50 55 60

Leu Asp Phe Trp Glu Met Val Trp His Asn Ser Gly Thr Asn Gly Val  
 65 70 75 80  
 Ile Val Met Leu Thr Asn Leu Tyr Glu Ala Gly Ser Glu Lys Cys Ser  
 85 90 95  
 Gln Tyr Trp Pro Asp Asn Lys Asp His Ala Leu Cys Leu Glu Gly Gly  
 100 105 110  
 Leu Arg Ile Ser Val Gln Lys Tyr Glu Thr Phe Glu Asp Leu Lys Val  
 115 120 125  
 His Leu Phe Arg Leu Asp Lys Pro Asn Gly Pro Pro Lys Tyr Ile His  
 130 135 140  
 His Phe Trp Val His Thr Trp Phe Asp Lys Thr His Pro Asp Ile Glu  
 145 150 155 160  
 Ser Ile Thr Gly Leu Ile Arg Cys Ile Asp Lys Val Pro Asn Asp Gly  
 165 170 175  
 Pro Met Phe Val His Cys Ser Ala Gly Val Gly Arg Thr Gly Thr Phe  
 180 185 190  
 Ile Ala Val Asp Gln Ile Leu Gln Val Pro Lys Asn Ile Leu Pro Lys  
 195 200 205  
 Thr Thr Asn Leu Glu Asp Ser Lys Asp Phe Ile Phe Asn Cys Val Asn  
 210 215 220  
 Ser Leu Arg Ser Gln Arg Met Lys Met Val Gln Asn Phe Glu Gln Phe  
 225 230 235 240  
 Lys Phe Leu Tyr Asp Val Val Asp Tyr Leu Ala Val Asp Gln Ile Leu  
 245 250 255  
 Gln Val Pro Lys Asn Ile Leu Pro Lys Thr Thr Asn Leu Glu Asp Ser  
 260 265 270  
 Lys Asp Phe Ile Phe Asn Cys Val Asn Ser Leu Arg Ser Gln Arg Met  
 275 280 285  
 Lys Met Val Gln Asn Phe Glu Gln Ile Lys Phe Leu Tyr Asp Val Val  
 290 295 300  
 Asp Tyr Leu  
 305

&lt;210&gt; 26

&lt;211&gt; 316

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 26

Gly Ile Thr Ala Asp Ser Ser Asn His Pro Asp Asn Lys His Lys Asn  
 1 5 10 15  
 Arg Tyr Ile Asn Ile Val Ala Tyr Asp His Ser Arg Val Lys Leu Ala  
 20 25 30  
 Gln Leu Ala Glu Lys Asp Gly Lys Leu Thr Asp Tyr Ile Asn Ala Asn  
 35 40 45  
 Tyr Val Asp Gly Tyr Asn Arg Pro Lys Ala Tyr Ile Ala Ala Gln Gly  
 50 55 60  
 Pro Leu Lys Ser Thr Ala Glu Asp Ile Thr Arg Met Ile Thr Glu His  
 65 70 75 80  
 Asn Val Glu Val Ile Val Met Ile Thr Asn Leu Val Gln Lys Gly Arg  
 85 90 95  
 Arg Lys Cys Asp Gln Tyr Trp Pro Ala Asp Gly Ser Gln Glu Tyr Gly  
 100 105 110  
 Asn Phe Leu Val Thr Gln Lys Ser Val Gln Val Leu Ala Tyr Tyr Thr

115	120	125
Val Ile Thr Leu Arg Asn Thr Lys Ile Lys Lys Gly Ser Gln Lys Gly		
130	135	140
Arg Pro Ser Gly Arg Val Thr Gln Tyr His Tyr Thr Gln Trp Pro		
145	150	155
Asp Met Gly Val Pro Glu Tyr Ser Leu Pro Val Leu Thr Phe Val Arg		160
	165	170
Lys Ala Ala Tyr Ala Lys Arg His Ala Val Gly Pro Val Val Val His		175
	180	185
Cys Ser Ala Gly Val Gly Arg Thr Gly Thr Tyr Ile Val Leu Asp Ser		190
	195	200
Met Leu Gln Gln Ile Gln His Glu Gly Thr Val Asn Ile Phe Gly Phe		205
	210	215
Leu Lys His Ile Arg Ser Gln Arg Asn Tyr Leu Val Gln Thr Glu Glu		220
225	230	235
Gln Tyr Val Phe Ile His Asp Thr Leu Val Glu Ala Ile Leu Ser Lys		240
	245	250
Gln Thr Glu Val Val Leu Asp Ser Met Leu Gln Gln Ile Gln His Glu		255
	260	265
Gly Thr Val Asn Ile Phe Gly Phe Leu Lys His Ile Arg Ser Gln Arg		270
	275	280
Asn Tyr Leu Val Gln Thr Glu Glu Gln Tyr Val Phe Ile His Asp Thr		285
	290	295
Leu Val Glu Ala Ile Leu Ser Lys Glu Thr Glu Val		300
305	310	315

&lt;210&gt; 27

&lt;211&gt; 294

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 27

Thr Ser Arg Phe Ile Ser Ala Asn Leu Pro Cys Asn Lys Phe Lys Asn	
1	5
Arg Leu Val Asn Ile Met Pro Tyr Glu Leu Thr Arg Val Cys Leu Gln	10
	20
Pro Ile Arg Gly Val Glu Gly Ser Asp Tyr Ile Asn Ala Ser Phe Leu	25
	30
Asp Gly Tyr Arg Gln Gln Lys Ala Tyr Ile Ala Thr Gln Gly Pro Leu	35
	40
Ala Glu Ser Thr Glu Asp Phe Trp Arg Met Leu Trp Glu His Asn Ser	45
65	55
Thr Ile Ile Val Met Leu Thr Lys Leu Arg Glu Met Gly Arg Glu Lys	60
	70
Cys His Gln Tyr Trp Pro Ala Glu Arg Ser Ala Arg Tyr Gln Tyr Phe	75
	80
Val Val Asp Pro Met Ala Glu Tyr Asn Met Pro Gln Tyr Ile Leu Phe	85
	90
Lys Val Thr Asp Ala Arg Asp Gly Gln Ser Arg Thr Ile Arg Gln Phe	95
	100
Gln Phe Thr Asp Trp Pro Glu Gln Gly Val Pro Lys Thr Gly Glu Gly	105
145	110
Phe Ile Asp Phe Ile Gly Gln Val His Lys Thr Lys Glu Gln Ile Gly	115
	120
	125
	130
	135
	140
	145
	150
	155
	160
	165
	170
	175

Gln Asp Gly Pro Ile Thr Val His Cys Ser Ala Gly Val Gly Arg Thr  
 180 185 190  
 Gly Val Phe Ile Thr Leu Ser Ile Val Leu Glu Arg Met Arg Tyr Glu  
 195 200 205  
 Gly Val Val Asp Met Phe Gln Thr Val Lys Thr Leu Arg Thr Gln Arg  
 210 215 220  
 Pro Ala Met Val Gln Thr Glu Asp Gln Tyr Gln Leu Cys Tyr Arg Ala  
 225 230 235 240  
 Ala Leu Glu Tyr Leu Thr Leu Ser Ile Val Leu Glu Arg Met Arg Tyr  
 245 250 255  
 Glu Gly Val Val Asp Met Phe Gln Thr Val Lys Thr Leu Arg Thr Gln  
 260 265 270  
 Arg Pro Ala Met Val Gln Thr Glu Asp Gln Tyr Gln Leu Cys Tyr Arg  
 275 280 285  
 Ala Ala Leu Glu Tyr Leu  
 290

<210> 28

<211> 281

<212> PRT

<213> Homo sapiens

<400> 28

Asn Asp Lys Met Arg Thr Gly Asn Leu Pro Ala Asn Met Lys Lys Asn  
 1 5 10 15  
 Arg Val Leu Gln Ile Ile Pro Tyr Glu Phe Asn Arg Val Ile Ile Pro  
 20 25 30  
 Val Lys Arg Gly Glu Asn Asp Lys Met Arg Thr Gly Asn Leu Pro Ala  
 35 40 45  
 Asn Met Lys Lys Asn Arg Val Leu Gln Ile Ile Pro Tyr Glu Phe Asn  
 50 55 60  
 Arg Val Ile Ile Pro Val Lys Arg Gly Glu Glu Asn Thr Asp Tyr Val  
 65 70 75 80  
 Asn Ala Ser Phe Ile Asp Gly Tyr Arg Gln Lys Asp Ser Tyr Ile Ala  
 85 90 95  
 Ser Gln Gly Pro Leu Leu His Thr Ile Glu Asp Phe Trp Arg Met Ile  
 100 105 110  
 Trp Glu Trp Lys Ser Cys Ser Ile Val Met Leu Thr Glu Leu Glu Glu  
 115 120 125  
 Arg Gly Gln Glu Lys Cys Ala Gln Tyr Trp Pro Ser Asp Gly Leu Val  
 130 135 140  
 Ser Tyr Gly Asp Ile Thr Val Glu Leu Lys Lys Glu Glu Glu Cys Glu  
 145 150 155 160  
 Ser Tyr Thr Val Leu Leu Val Thr Asn Thr Arg Glu Asn Lys Ser Arg  
 165 170 175  
 Gln Ile Arg Gln Phe His Phe His Gly Trp Pro Glu Val Gly Ile Pro  
 180 185 190  
 Ser Asp Gly Lys Gly Met Ile Ser Ile Ile Ala Ala Val Gln Lys Gln  
 195 200 205  
 Gln Gln Gln Ser Gly Asn His Pro Ile Thr Val His Cys Ser Ala Gly  
 210 215 220  
 Ala Gly Arg Thr Gly Thr Phe Cys Ala Leu Ser Thr Val Leu Glu Arg  
 225 230 235 240  
 Val Lys Ala Glu Gly Ile Leu Asp Val Phe Gln Thr Val Lys Ser Leu



245 250 255  
 Arg Leu Gln Arg Pro His Met Val Gln Thr Leu Glu Gln Tyr Glu Phe  
 260 265 270  
 Cys Tyr Lys Val Val Gln Glu Tyr Ile  
 275 280

<210> 29

<211> 298

<212> PRT

<213> Homo sapiens

<400> 29

Lys Glu Asn Met Arg Thr Gly Asn Leu Pro Ala Asn Met Lys Lys Ala  
 5 10 15  
 Arg Val Ile Gln Ile Ile Pro Tyr Asp Phe Asn Arg Val Ile Leu Ser  
 20 25 30  
 Met Lys Arg Gly Gln Glu Tyr Thr Asp Tyr Ile Asn Ala Ser Phe Ile  
 35 40 45  
 Asp Gly Tyr Arg Gln Lys Asp Tyr Phe Ile Ala Thr Gln Gly Pro Leu  
 50 55 60  
 Ala His Thr Val Glu Asp Phe Trp Arg Met Ile Trp Glu Trp Lys Ser  
 65 70 75 80  
 His Thr Ile Val Met Leu Thr Glu Val Gln Glu Arg Glu Gln Asp Lys  
 85 90 95  
 Cys Tyr Gln Tyr Trp Pro Thr Glu Gly Ser Val Thr His Gly Glu Ile  
 100 105 110  
 Thr Ile Glu Ile Lys Asn Asp Thr Leu Ser Glu Ala Ile Ser Ile Phe  
 115 120 125  
 Leu Val Thr Leu Asn Gln Pro Gln Ala Arg Gln Glu Glu Gln Val Arg  
 130 135 140  
 Val Val Arg Gln Phe His Phe His Gly Trp Pro Glu Ile Gly Ile Pro  
 145 150 155 160  
 Ala Glu Gly Lys Gly Met Ile Asp Leu Ile Ala Ala Val Gln Lys Gln  
 165 170 175  
 Gln Gln Gln Thr Gly Asn His Pro Ile Thr Val His Cys Ser Ala Gly  
 180 185 190  
 Ala Gly Arg Thr Gly Thr Phe Ile Ala Leu Ser Asn Ile Leu Glu Arg  
 195 200 205  
 Val Lys Ala Glu Gly Leu Leu Asp Val Phe Gln Ala Val Lys Ser Leu  
 210 215 220  
 Arg Leu Gln Arg Pro His Met Val Gln Thr Leu Glu Gln Tyr Glu Phe  
 225 230 235 240  
 Cys Tyr Lys Val Val Gln Asp Phe Ile Ala Leu Ser Asn Ile Leu Glu  
 245 250 255  
 Arg Val Lys Ala Glu Gly Leu Leu Asp Val Phe Gln Ala Val Lys Ser  
 260 265 270  
 Leu Arg Leu Gln Arg Pro His Met Val Gln Thr Leu Glu Gln Tyr Glu  
 275 280 285  
 Phe Cys Tyr Lys Val Val Gln Asp Phe Ile  
 290 295

<210> 30

<211> 301

<212> PRT

\*213\* H. m. capiens

\*400\* 30

Val	Glu	Asp	Cys	Ser	Ile	Ala	Leu	Leu	Pro	Arg	Asn	His	Glu	Lys	Asn
1			5						10					15	
Arg	Cys	Met	Asp	Ile	Leu	Pro	Pro	Asp	Arg	Cys	Leu	Pro	Phe	Leu	Ile
		20						25					30		
Thr	Ile	Asp	Gly	Glu	Ser	Ser	Asn	Tyr	Ile	Asn	Ala	Ala	Leu	Met	Asp
		35					40					45			
Ser	Tyr	Lys	Gln	Pro	Ser	Ala	Phe	Ile	Val	Thr	Gln	His	Pro	Leu	Pro
	50					55					60				
Asn	Thr	Val	Lys	Asp	Phe	Tyr	Arg	Leu	Val	Leu	Asp	Tyr	His	Cys	Thr
65					70					75				80	
Ser	Val	Val	Met	Leu	Asn	Asp	Val	Asp	Pro	Ala	Gln	Leu	Cys	Pro	Gln
			85					90					95		
Tyr	Trp	Pro	Glu	Asn	Gly	Val	His	Arg	His	Gly	Pro	Ile	Gln	Val	Glu
		100						105					110		
Phe	Val	Ser	Ala	Asp	Leu	Glu	Glu	Asp	Ile	Ile	Ser	Phe	Arg	Ile	Tyr
	115					120						125			
Asn	Ala	Ala	Arg	Pro	Gln	Asp	Gly	Tyr	Arg	Met	Val	Gln	Gln	Phe	Gln
	130				135						140				
Phe	Leu	Gly	Tyr	Pro	Met	Tyr	Arg	Asp	Thr	Pro	Val	Ser	Lys	Arg	Ser
145					150					155				160	
Phe	Leu	Lys	Leu	Ile	Arg	Gln	Val	Asp	Lys	Trp	Gln	Glu	Glu	Tyr	Asn
			165					170						175	
Gly	Gly	Glu	Gly	Pro	Thr	Val	Val	His	Cys	Leu	Asn	Gly	Gly	Gly	Arg
		180						185					190		
Ser	Gly	Thr	Phe	Cys	Ala	Ile	Ser	Ile	Val	Cys	Glu	Met	Leu	Arg	His
	195					200						205			
Gln	Arg	Thr	Val	Asp	Val	Phe	His	Ala	Val	Lys	Thr	Leu	Arg	Asn	Asn
	210					215					220				
Lys	Pro	Asn	Met	Val	Asp	Leu	Leu	Asp	Gln	Tyr	Lys	Phe	Cys	Tyr	Glu
225					230					235				240	
Val	Ala	Leu	Glu	Tyr	Leu	Asn	Ser	Gly	Ala	Ile	Ser	Ile	Val	Cys	Glu
			245					250						255	
Met	Leu	Arg	His	Gly	Arg	Thr	Val	Asp	Val	Phe	His	Ala	Val	Lys	Thr
		260				265							270		
Leu	Arg	Asn	Asn	Lys	Pro	Asn	Met	Val	Asp	Leu	Leu	Asp	Gln	Tyr	Lys
		275				280						285			
Phe	Cys	Tyr	Glu	Val	Ala	Leu	Glu	Tyr	Leu	Asn	Ser	Gly			
	290					295					300				

\*210\* 31

\*211\* 333

\*212\* PKT

\*213\* Mus musculus

\*400\* 31

Trp	Arg	Thr	Glu	His	Ile	Gly	Asn	Gln	Glu	Glu	Asn	Lys	Lys	Lys	Asn
1				5					10					15	
Arg	Asn	Ser	Asn	Val	Val	Pro	Tyr	Asp	Phe	Asn	Arg	Val	Pro	Leu	Lys
		20						25					30		
His	Glu	Leu	Glu	Met	Ser	Lys	Glu	Ser	Glu	Pro	Glu	Ser	Asp	Glu	Ser
		35					40					45			

Ser Asp Asp Asp Ser Asp Ser Glu Glu Thr Ser Lys Tyr Ile Asn Ala  
 50 60  
 Ser Phe Val Met Ser Tyr Trp Lys Pro Glu Met Met Ile Ala Ala Gln  
 65 70 75 80  
 Gly Pro Leu Lys Glu Thr Ile Gly Asp Phe Trp Gln Met Ile Phe Gln  
 85 90 95  
 Arg Lys Val Lys Val Ile Val Met Leu Thr Glu Leu Val Asn Gly Asp  
 100 105 110  
 Gln Glu Val Cys Ala Gln Tyr Trp Gly Glu Gly Lys Gln Thr Tyr Gly  
 115 120 125  
 Asp Met Glu Val Glu Met Lys Asp Thr Asn Arg Ala Ser Ala Tyr Thr  
 130 135 140  
 Leu Phe Glu Leu Arg His Ser Lys Arg Lys Glu Pro Arg Thr Val Tyr  
 145 150 155 160  
 Gln Tyr Gln Cys Thr Thr Trp Lys Gly Glu Glu Leu Pro Ala Glu Pro  
 165 170 175  
 Lys Asp Leu Val Ser Met Ile Gln Asp Leu Lys Gln Lys Leu Pro Lys  
 180 185 190  
 Ala Ser Pro Glu Gly Met Lys Tyr His Lys His Ala Ser Ile Leu Val  
 195 200 205  
 His Cys Arg Asp Gly Ser Gln Gln Thr Gly Leu Phe Cys Ala Leu Phe  
 210 215 220  
 Asn Leu Leu Glu Ser Ala Glu Thr Glu Asp Val Val Asp Val Phe Gln  
 225 230 235 240  
 Val Val Lys Ser Leu Arg Lys Ala Arg Pro Gly Val Val Cys Ser Tyr  
 245 250 255  
 Glu Gln Tyr Gln Phe Leu Tyr Asp Ile Ile Ala Ser Ile Tyr Pro Ala  
 260 265 270  
 Gln Asn Gly Gln Val Ala Leu Phe Asn Leu Leu Glu Ser Ala Glu Thr  
 275 280 285  
 Glu Asp Val Val Asp Val Phe Gln Val Val Lys Ser Leu Arg Lys Ala  
 290 295 300  
 Arg Pro Gly Val Val Cys Ser Tyr Glu Gln Tyr Gln Phe Leu Tyr Asp  
 305 310 315 320  
 Ile Ile Ala Ser Ile Tyr Pro Ala Gln Asn Gly Gln Val  
 325 330

&lt;210&gt; 32

&lt;211&gt; 295

&lt;212&gt; PRT

<213> *Drosophila melanogaster*

&lt;400&gt; 32

Ser Lys Ser Cys Ser Val Gly Glu Asn Glu Glu Asn Asn Met Lys Asn  
 1 10 15  
 Arg Ser Gln Glu Ile Ile Pro Tyr Asp Arg Asn Arg Val Ile Leu Thr  
 20 25 30  
 Pro Leu Pro Met Arg Glu Asn Ser Thr Tyr Ile Asn Ala Ser Phe Ile  
 35 40 45  
 Glu Gly Tyr Asp Asn Ser Glu Thr Phe Ile Ile Ala Gln Asp Pro Phe  
 50 55 60  
 Glu Asn Thr Ile Gly Asp Phe Tyr Arg Met Ile Ser Glu Gln Ser Val  
 65 70 75 80  
 Thr Thr Leu Val Met Ile Ser Glu Ile Gly Asp Gly Pro Arg Lys Cys

85 90 95  
 Pro Arg Tyr Trp Ala Asp Asp Gln Val Gln Tyr Asp His Ile Leu Val  
 100 105 110  
 Lys Tyr Val His Ser Glu Ser Cys Pro Tyr Tyr Thr Phe Phe Tyr Val  
 115 120 125  
 Thr Asn Cys Lys Ile Asp Asp Thr Leu Lys Val Thr Gln Phe Gln Tyr  
 130 135 140  
 Asn Gly Trp Pro Thr Val Asp Gly Glu Val Pro Glu Val Cys Arg Gly  
 145 150 155 160  
 Ile Ile Glu Leu Val Asp Gln Ala Tyr Asn His Tyr Lys Asn Asn Lys  
 165 170 175  
 Asn Ser Gly Cys Arg Ser Pro Leu Thr Val His Cys Ser Leu Gly Thr  
 180 185 190  
 Asp Arg Ser Ser Ile Phe Val Ala Met Cys Ile Leu Val Gln His Leu  
 195 200 205  
 Arg Leu Glu Lys Cys Val Asp Ile Cys Ala Thr Thr Arg Lys Leu Arg  
 210 215 220  
 Ser Gln Arg Thr Gly Leu Ile Asn Ser Tyr Ala Gln Tyr Gln Phe Leu  
 225 230 235 240  
 His Arg Ala Ile Ile Asn Tyr Ala Met Cys Ile Leu Val Gln His Leu  
 245 250 255  
 Arg Leu Glu Lys Cys Val Asp Ile Cys Ala Thr Thr Arg Lys Leu Arg  
 260 265 270  
 Ser Gln Arg Thr Gly Leu Ile Asn Ser Tyr Ala Gln Tyr Glu Phe Leu  
 275 280 285  
 His Arg Ala Ile Ile Asn Tyr  
 290 295

<210> 33

<211> 308

<212> PRT

<213> Homo sapiens

<400> 33

Gln Ser Asp Tyr Ser Ala Ala Leu Lys Gln Cys Asn Arg Glu Lys Asn  
 1 5 10 15  
 Arg Thr Ser Ser Ile Ile Pro Val Glu Arg Ser Arg Val Gly Ile Ser  
 20 25 30  
 Ser Leu Ser Gly Glu Gly Thr Asp Tyr Ile Asn Ala Ser Tyr Ile Met  
 35 40 45  
 Gly Tyr Tyr Gln Ser Asn Glu Phe Ile Ile Thr Gln His Pro Leu Leu  
 50 55 60  
 His Thr Ile Lys Asp Phe Trp Arg Met Ile Trp Asp His Asn Ala Gln  
 65 70 75 80  
 Leu Val Val Met Ile Pro Asp Gly Gln Asn Met Ala Gln Asp Glu Phe  
 85 90 95  
 Val Tyr Trp Pro Asn Lys Asp Glu Pro Ile Asn Cys Glu Ser Phe Lys  
 100 105 110  
 Val Thr Leu Met Ala Glu Gln His Lys Cys Leu Ser Asn Gln Glu Lys  
 115 120 125  
 Leu Ile Ile Ile Ile Leu Glu Ala Thr Gln Asp Asp Tyr Val Leu Glu  
 130 135 140  
 Val Arg His Phe Gln Cys Pro Lys Trp Pro Asn Ile Asp Ser Pro Ile  
 145 150 155 160

Ser Lys Thr Phe Glu Leu Ile Ser Val Ile Lys Glu Glu Ala Ala Asn  
 165 170 175  
 Arg Asp Gly Pro Met Ile Val His Asp Glu His Gly Gly Val Thr Ala  
 180 185 190  
 Gly Thr Phe Cys Ala Leu Thr Thr Leu Met His Gln Leu Glu Lys Glu  
 195 200 205  
 Asn Ser Val Asp Val Tyr Gln Val Ala Lys Met Ile Asn Leu Met Arg  
 210 215 220  
 Pro Gly Val Phe Ala Asp Ile Glu Gln Tyr Gln Phe Leu Tyr Lys Val  
 225 230 235 240  
 Ile Leu Ser Leu Val Ser Thr Arg Gln Glu Glu Asn Ala Leu Thr Thr  
 245 250 255  
 Leu Met His Gln Leu Glu Lys Glu Asn Ser Val Asp Val Tyr Gln Val  
 260 265 270  
 Ala Lys Met Ile Asn Leu Met Arg Pro Gly Val Phe Ala Asp Ile Glu  
 275 280 285  
 Gln Tyr Gln Phe Leu Tyr Lys Val Ile Leu Ser Leu Val Ser Thr Arg  
 290 295 300  
 Gln Glu Glu Asn  
 305

&lt;210&gt; 34

&lt;211&gt; 308

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 34

Val Glu Cys Phe Ser Ala Gln Lys Glu Cys Asn Lys Glu Lys Asn Arg  
 1 5 10 15  
 Asn Ser Ser Val Val Pro Ser Glu Arg Ala Arg Val Gly Leu Ala Pro  
 20 25 30  
 Leu Pro Gly Met Lys Gly Thr Asp Tyr Ile Asn Ala Ser Tyr Ile Met  
 35 40 45  
 Gly Tyr Tyr Arg Ser Asn Glu Phe Ile Ile Thr Gln His Pro Leu Pro  
 50 55 60  
 His Thr Thr Lys Asp Phe Trp Arg Met Ile Trp Asp His Asn Ala Gln  
 65 70 75 80  
 Ile Ile Val Met Leu Pro Asp Asn Gln Ser Leu Ala Glu Asp Glu Ile  
 85 90 95  
 Val Tyr Trp Pro Ser Arg Glu Glu Ser Met Asn Cys Glu Ala Phe Thr  
 100 105 110  
 Val Thr Leu Ile Ser Lys Asp Arg Leu Cys Leu Ser Asn Glu Glu Gln  
 115 120 125  
 Ile Ile Ile Phe Ile Leu Glu Ala Thr Gln Asp Asp Tyr Val Leu Glu  
 130 135 140  
 Val Arg His Phe Gln Cys Pro Lys Trp Pro Asn Pro Asp Ala Pro Ile  
 145 150 155 160  
 Ser Ser Thr Phe Glu Leu Ile Asn Val Ile Lys Glu Glu Ala Leu Thr  
 165 170 175  
 Arg Asp Gly Pro Thr Ile Val His Asp Gln Tyr Gly Ala Val Ser Ala  
 180 185 190  
 Gly Met Leu Cys Ala Leu Thr Thr Leu Ser Gln Gln Leu Glu Asn Glu  
 195 200 205  
 Asn Ala Val Asp Val Phe Gln Val Ala Lys Met Ile Asn Leu Met Arg

210 215 220  
 Pro Gly Val Phe Thr Asp Ile Glu Gln Tyr Gln Phe Ile Tyr Lys Ala  
 225 230 235 240  
 Met Leu Ser Leu Val Ser Thr Lys Gln Asn Gly Asn Ala Leu Thr Thr  
 245 250 255  
 Leu Ser Gln Gln Leu Glu Asn Glu Asn Ala Val Asp Val Phe Gln Val  
 260 265 270  
 Ala Lys Met Ile Asn Leu Met Arg Pro Gly Val Phe Thr Asp Ile Glu  
 275 280 285  
 Gln Tyr Gln Phe Ile Tyr Lys Ala Met Leu Ser Leu Val Ser Thr Lys  
 290 295 300  
 Glu Asn Gly Asn  
 305

<210> 35

<211> 335

<212> ERT

<213> *Drosophila melanogaster*

<400> 35

Glu Thr Asn Leu Met Ala Glu Gln Val Glu Glu Leu Lys Asn Cys Thr  
 1 5 10 15  
 Pro Tyr Leu Glu Gln Gln Tyr Lys Asn Ile Ile Gln Phe Gln Pro Lys  
 20 25 30  
 Asp Ile His Ile Ala Ser Ala Met Lys Gln Val Asn Ser Ile Lys Asn  
 35 40 45  
 Arg Gly Ala Ile Phe Pro Ile Glu Gly Ser Arg Val His Leu Thr Pro  
 50 55 60  
 Lys Pro Gly Glu Asp Gly Ser Asp Tyr Ile Asn Ala Ser Trp Leu His  
 65 70 75 80  
 Gly Phe Arg Arg Leu Arg Asp Phe Ile Val Thr Gln His Pro Met Ala  
 85 90 95  
 His Thr Ile Lys Asp Phe Trp Gln Met Val Trp Asp His Asn Ala Gln  
 100 105 110  
 Thr Val Val Leu Leu Ser Ser Leu Asp Asp Ile Asn Phe Ala Gln Phe  
 115 120 125  
 Trp Pro Asp Glu Ala Thr Pro Ile Glu Ser Asp His Tyr Arg Val Lys  
 130 135 140  
 Phe Leu Asn Lys Thr Asn Lys Ser Asp Tyr Val Ser Phe Val Ile Gln  
 145 150 155 160  
 Ser Ile Gln Asp Asp Tyr Glu Leu Thr Val Lys Met Leu His Cys Pro  
 165 170 175  
 Ser Trp Pro Glu Met Ser Asn Pro Asn Ser Ile Tyr Asp Phe Ile Val  
 180 185 190  
 Asp Val His Glu Arg Cys Asn Asp Tyr Arg Asn Gly Pro Ile Val Ile  
 195 200 205  
 Val Asp Arg Tyr Gly Gly Ala Gln Ala Cys Thr Ile Cys Ala Ile Ser  
 210 215 220  
 Ser Leu Ala Ile Glu Met Glu Tyr Cys Ser Thr Ala Asn Val Tyr Gln  
 225 230 235 240  
 Tyr Ala Lys Leu Tyr His Asn Lys Arg Pro Gly Val Trp Thr Ser Ser  
 245 250 255  
 Glu Asp Ile Arg Val Ile Tyr Asn Ile Leu Ser Phe Leu Pro Gly Asn  
 260 265 270

Leu Asn Leu Leu Lys Arg Ala Ile Ser Ser Leu Ala Ile Gln Met Gln  
 275 280 285  
 Tyr Cys Ser Thr Ala Asn Val Tyr Gln Tyr Ala Lys Leu Tyr His Asn  
 290 295 300  
 Lys Arg Pro Gly Val Trp Ser Ser Glu Asp Ile Arg Val Ile Tyr  
 305 310 315 320  
 Asn Ile Leu Ser Phe Leu Pro Gly Asn Leu Asn Leu Leu Lys Arg  
 325 330 335

<210> 36

<211> 237

<212> PRT

<213> Yersinia sp.

<400> 36

Thr Asn Asp Pro Arg Tyr Leu Gln Ala Cys Gly Gly Glu Lys Ile Leu  
 : 5 10 15  
 Asn Arg Phe Arg Asp Ile Gln Cys Cys Arg Gln Thr Ala Val Arg Ala  
 20 25 30  
 Asp Asn Tyr Ile Gln Val Gly Asn Thr Arg Thr Ile Ala Cys Gln Tyr  
 35 40 45  
 Pro Leu Gln Ser Gln Leu Glu Ser His Phe Arg Met Leu Ala Glu Asn  
 50 55 60  
 Arg Thr Pro Val Leu Ala Val Leu Ala Ser Ser Ser Glu Ile Ala Asn  
 65 70 75 80  
 Gln Arg Phe Gly Met Pro Asp Tyr Phe Arg Gln Ser Gly Thr Tyr Gly  
 85 90 95  
 Ser Ile Thr Val Glu Ser Lys Met Thr Gln Gln Val Gly Leu Gly Asp  
 100 105 110  
 Gly Ile Asn Met Tyr Thr Leu Thr Ile Arg Glu Ala Gly Gln Lys Thr  
 115 120 125  
 Ile Ser Val Pro Val Val His Val Gly Asn Trp Pro Asp Gln Thr Ala  
 130 135 140  
 Val Ser Ser Glu Val Thr Lys Ala Leu Ala Ser Leu Val Asp Gln Thr  
 145 150 155 160  
 Ala Glu Thr Lys Arg Asn Met Tyr Glu Ser Lys Gly Ser Ser Ala Val  
 165 170 175  
 Ala Asp Asp Ser Lys Leu Arg Pro Val Ile His Cys Arg Ala Gly Val  
 180 185 190  
 Gly Arg Thr Ala Gln Leu Ile Gly Ala Met Cys Met Asn Asp Ser Arg  
 195 200 205  
 Asn Ser Gln Leu Ser Val Glu Asp Met Val Ser Gln Met Arg Val Gln  
 210 215 220  
 Arg Asn Gly Met Val Gln Lys Asp Glu Gln Leu Asp Val Leu Ile Lys  
 225 230 235 240  
 Leu Ala Glu Gly Ala Met Cys Met Asn Asp Ser Arg Asn Ser Gln Leu  
 245 250 255  
 Ser Val Glu Asp Met Val Ser Gln Met Arg Val Gln Arg Asn Gly Met  
 260 265 270  
 Val Gln Lys Asp Glu Gln Leu Asp Val Leu Ile Lys Leu Ala Glu  
 275 280 285

<410> 37

<411> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> Fluorescently-labeled phosphopeptides derived from amino acids 1170-1176 of the EGF receptor sequence.

<221> PHOSPHORYLATION

<222> (4)...(4)

<400> 37

Asn Ala Glu Tyr Leu Arg Val

1 5

<210> 38

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Preferred substrate for PTB1B, corresponding to residues 988-993 of human EGF receptor.

<221> PHOSPHORYLATION

<222> (5)...(5)

<400> 38

Arg Ala Asp Glu Tyr Leu

1 5

<210> 39

<211> 11

<212> PRT

<213> Artificial Sequence

<220>

<223> Substrate for PTBs synthesized from residues 1142-1152 of human insulin receptor.

<221> PHOSPHORYLATION

<222> (5)...(5)

<400> 39

His Arg Asp Ile Tyr Glu Thr Asp Tyr Tyr Arg

1 5 10

<210> 40

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> Substrate for PTBs synthesized from residues



600-500 of p16<sup>lck</sup>, the src-like lymphocyte specific protein tyrosine kinase that is a physiological substrate for CD45.

#### \*221\* PHOSPHORYLATION

\*222\* (6)...(6)

\*400\* 40

Ala Thr Gln Gly Gln Tyr Gln Pro Gln Pro  
1 5 10